

The Variant Effect Predictor predicts the functional effects of genomic variants <http://www.ensembl.org/vep>

824 commits 10 branches 52 releases 10 contributors

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Laurent ens-igil committed on GitHub Merge pull request #73 from MaxUlysse/patch-1	
docker	sorted tools and removed duplicates
examples	add example VCF inputs
modules/Bio/EnsEMBL/VEP	escape spaces in symbol names for VCF
t	bypass get_raw_end method for non-SVs, SVLEN in INFO can mess up coords

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Use Git or checkout with SVN using the web URL.

<https://github.com/Ensembl/ensembl-vep.git>

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